

## SEQUENCE LISTING

<110> Hong Zhang  
Andrew T. Watt

<120> ANTISENSE MODULATION OF CASPASE 7 EXPRESSION

<130> RTS-0201

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<213> Artificial Sequence

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<213> Homo sapiens

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[illegible]

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<213> Artificial Sequence

<223> PCR Primer

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<210> 6

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<212> DNA

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<223> PCR Probe

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<211> 26

RTS-0201

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;223&gt; PCR Primer

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26

&lt;210&gt; 9

&lt;211&gt; 21

&lt;212&gt; DNA

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&lt;223&gt; PCR Probe

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21

&lt;210&gt; 10

&lt;211&gt; 2006

&lt;212&gt; DNA

&lt;213&gt; Mus musculus

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (474)...(1496)

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Pro Ser Thr Leu Ala Glu Asp Gly Arg Cys Arg Gly Leu Leu Ala Ala	
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Ala Val Gly Thr Met Thr Asp Asp Gln Asp Cys Ala Ala Glu Leu Glu	
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Lys Val Asp Ser Ser Ser Glu Asp Gly Val Asp Ala Lys Pro Asp Arg	
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Lys Ala Ser Glu Glu Asp His Ser Asn Ser Ala Cys Phe Ala Cys Val	
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Thr Pro Ile Lys Asp Leu Thr Ala His Phe Arg Gly Asp Arg Cys Lys	
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Val Ala Arg His Phe Glu Ser Gln Ser Asp Asp Pro Arg Phe Asn Glu	
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Phe Ser Arg  
340

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<223> PCR Primer

<400> 11

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<210> 12

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<213> Homo sapiens

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ggatttgtgc ttcttatgtt acccag atg gca gat gat cag ggc tgt att gaa 413  
Met Ala Asp Asp Gln Gly Cys Ile Glu  
1 5

gag cag ggg gtt gag gat tca gca aat gaa gat tca gtg gat gct aag 461  
Glu Gln Gly Val Glu Asp Ser Ala Asn Glu Asp Ser Val Asp Ala Lys  
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cca gac cgg tcc tcg ttt gta ccg tcc ctc ttc agt aag aag aag aaa 509  
Pro Asp Arg Ser Ser Phe Val Pro Ser Leu Phe Ser Lys Lys Lys Lys  
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aat gtc acc atg cga tcc atc aag acc acc cgg gac cga gtg cct aca 557  
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45 50 55

tat cag tac aac atg aat ttt gaa aag ctg ggc aaa tgc atc ata ata 605  
Tyr Gln Tyr Asn Met Asn Phe Glu Lys Leu Gly Lys Cys Ile Ile Ile  
60 65 70

aac aac aag aac ttt gat aaa gtg aca ggt atg ggc gtt cga aac gga 653  
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Leu Leu Lys Lys Ala Ser Glu Glu Asp His Thr Asn Ala Ala Cys Phe	
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Asp Gly Val Thr Pro Ile Lys Asp Leu Thr Ala His Phe Arg Gly Asp	
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Arg Cys Lys Thr Leu Leu Glu Lys Pro Lys Leu Phe Phe Ile Gln Ala	
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Cys Arg Gly Thr Glu Leu Asp Asp Gly Ile Gln Ala Asp Ser Gly Pro	
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tgccctcaca tcttcacca gcaccttact gccaggcct atctggaagc cacctacca 2228

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RTS-0201

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<221> CDS

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 Met Asp Cys  
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gtt ggt tgg cct cca ggc agg aag tgg cac ttg gaa aag aac acc agc 285  
 Val Gly Trp Pro Pro Gly Arg Lys Trp His Leu Glu Lys Asn Thr Ser  
 5 10 15

tgc ggt ggt agc agt ggg att tgt gct tct tat gtt acc cag atg gca 333  
 Cys Gly Gly Ser Ser Gly Ile Cys Ala Ser Tyr Val Thr Gln Met Ala  
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RTS-0201 "PATENT"

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40 45 50	
 gaa gat tca gtg gat gct aag cca gac cgg tcc tcg ttt gta ccg tcc	429
Glu Asp Ser Val Asp Ala Lys Pro Asp Arg Ser Ser Phe Val Pro Ser	
55 60 65	
 ctc ttc agt aag aag aag aaa aat gtc acc atg cga tcc atc aag acc	477
Leu Phe Ser Lys Lys Lys Lys Asn Val Thr Met Arg Ser Ile Lys Thr	
70 75 80	
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Thr Arg Asp Arg Val Pro Thr Tyr Gln Tyr Asn Met Asn Phe Glu Lys	
85 90 95	
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Leu Gly Lys Cys Ile Ile Ile Asn Asn Lys Asn Phe Asp Lys Val Thr	
100 105 110 115	
 ggg atg ggc gtt cga aac gga aca gac aaa gat gcc gag gcg ctc ttc	621
Gly Met Gly Val Arg Asn Gly Thr Asp Lys Asp Ala Glu Ala Leu Phe	
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 aag tgc ttc cga agc ctg ggt ttt gac gtg att gtc tat aat gac tgc	669
Lys Cys Phe Arg Ser Leu Gly Phe Asp Val Ile Val Tyr Asn Asp Cys	
135 140 145	
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Ser Cys Ala Lys Met Gln Asp Leu Leu Lys Lys Ala Ser Glu Glu Asp	
150 155 160	
 cat aca aat gcc gcc tgc ttc gcc tgc atc ctc tta agc cat gga gaa	765
His Thr Asn Ala Ala Cys Phe Ala Cys Ile Leu Leu Ser His Gly Glu	
165 170 175	
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Glu Asn Val Ile Tyr Gly Lys Asp Gly Val Thr Pro Ile Lys Asp Leu	
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Thr Ala His Phe Arg Gly Asp Arg Cys Lys Thr Leu Leu Glu Lys Pro	
200 205 210	

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75 80 85

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&lt;211&gt; 20

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&lt;223&gt; Antisense Oligonucleotide

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RTS-0201

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RTS-0201

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Ile Ile Ser Ser Ile Leu Leu Lys Lys Lys Arg Asn Ala Ser Ala Gly	
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Asp Lys Ala Thr Gly Met Asp Val Arg Asn Gly Thr Asp Lys Asp Ala	
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Gly Ala Leu Phe Lys Cys Phe Gln Asn Leu Gly Phe Glu Val Thr Val	
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His Asn Asp Cys Ser Cys Ala Lys Met Gln Asp Leu Leu Arg Lys Ala	
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Ser Glu Glu Asp His Ser Asn Ser Ala Cys Phe Ala Cys Val Leu Leu	
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Ser His Gly Glu Glu Asp Leu Ile Tyr Gly Lys Asp Gly Val Thr Pro	
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Ile Lys Asp Leu Thr Ala His Phe Arg Gly Asp Arg Cys Lys Thr Leu	
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Leu Asp Asp Gly Ile Gln Ala Asp Ser Gly Pro Ile Asn Asp Ile Asp	
195 200 205	
gct aat ccc cgc aac aag atc ccg gtg gaa gcc gac ttc ctc ttt gct	855
Ala Asn Pro Arg Asn Lys Ile Pro Val Glu Ala Asp Phe Leu Phe Ala	
210 215 220	
tac tcc acg gtt cca ggt tat tac tca tgg agg aac cca ggg aaa ggc	903
Tyr Ser Thr Val Pro Gly Tyr Tyr Ser Trp Arg Asn Pro Gly Lys Gly	
225 230 235	
tcc tgg ttt gtg cag gcc ctc tgc tcc atc ctg aat gag cat ggc aag	951
Ser Trp Phe Val Gln Ala Leu Cys Ser Ile Leu Asn Glu His Gly Lys	
240 245 250	
gac ctc gag atc atg cag atc ctg acc agg gtg aac gac agg gtg gcc	999
Asp Leu Glu Ile Met Gln Ile Leu Thr Arg Val Asn Asp Arg Val Ala	
255 260 265 270	
agg cac ttc gag tcc cag tct gat gat cca cgc ttc aac gag aag aag	1047
Arg His Phe Glu Ser Gln Ser Asp Asp Pro Arg Phe Asn Glu Lys Lys	
275 280 285	

cag atc ccg tgt atg gtg tcc atg ctc acc aaa gag ctg tac ttc agc	1095
Gln Ile Pro Cys Met Val Ser Met Leu Thr Lys Glu Leu Tyr Phe Ser	
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cgt tga ccacccttca gctgagaacc tgccgccgtt cgttgatgaa tccagttttt	1151
Arg	
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ggaaagtcta gatttttttt tttttgttta ataactttgt tcatctgatg acttcatgct	1271
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aatgtgaatt aaataaccat gtcctctggc gtgctacaat gtattcattc actattcact	2051
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agcagtttct ttccaaggta tcatttttctt tggaattcgt tatacattcc tgtttttccc 2291  
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tccagcgaag acggagttga cgccaagcca gaccgctcct ctatcatctc ctctattctc 180  
ttgaagaaga agagaaatgc ctctgggggc cccgtcagga ccggccggtg cgcgggtgcc 240  
cactggtggg gaccgcaggg aggtggcaag tgaggagag gtgcatgggt ggagagaaga 300  
ggggcgctcg caaagcgagg tttgtggagg tccggaatgg gacggagaaa gatgcagggg 360  
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<400> 167

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<213> Artificial Sequence

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